WO 2004/015067

SEO ID NO: 1 (Human CatSper 3 cDNA)

PCT/US2003/024432 10/5-234-79 DT01 Rec'd PCT/PTC 0 4 FFR 2005

ATGAGGGATA ATGAAAAGGC CTGGTGGCAG CAATGGACCT CCCATACAGG 0050 CCTCGAGGGG TGGGGCGGGA CTCAGGAGGA CCGTATGGGG TTTGGAGGGG 0100 CAGTAGCTGC ACTGAGGGGC CGCCCCTCTC CCCTGCAGAG TACCATTCAC 0150 GAGTCCTACG GTCGGCCAGA GGAGCAAGTG CTCATCAACC GCCAGGAAAT 0200 CACGAACAAA GCGGACGCCT GGGACATGCA GGAGTTCATC ACTCACATGT 0250 ACATCAAGCA GCTGCTCCGA CACCCCGCCT TCCAACTGCT GCTGGCCCTG 0300 0350 CTGCTGGTGA TCAATGCCAT CACCATCGCT CTCCGTACCA ACTCCTACCT GGACCAGAAA CACTATGAGT TGTTCTCTAC CATAGATGAC ATTGTGCTGA 0400 CCATCCTTCT TTGTGAGGTT CTCCTTGGCT GGCTCAATGG CTTCTGGATT 0450 TTCTGGAAGG ACGGCTGGAA CATCCTCAAC TTCATTATCG TCTTTATCTT 0500 GCTCTTGCGG TTCTTCATTA ATGAAATCAA TATTCCCTCC ATCAACTACA 0550 CTCTCAGGGC GCTTCGTCTG GTGCATGTGT GCATGGCGGT GGAGCCCCTC 0600 GCCCGGATCA TCCGCGTCAT CCTGCAGTCG GTGCCTGACA TGGCCAATAT 0650 CATGGTCCTC ATCCTCTTCT TCATGCTGGT TTTTTCCGTG TTTGGAGTAA 0700 CACTCTTTGG TGCATTCGTG CCCAAGCATT TCCAGAACAT ACAGGTTGCG 0750 CTGTACACCC TCTTCATCTG CATCACCCAG GACGGCTGGG TGGACATCTA 0800 CAGTGACTTC CAGACAGAGA AGAGGGAATA TGCAATGGAG ATTGGGGGTG 0850 CCATCTACTT TACCATCTTC ATCACCATCG GTGCCTTCAT TGGCATCAAC 0900 CTGTTCGTCA TCGTGGTGAC CACCAACCTG GAGCAAATGA TGAAGGCAGG 0950 AGAGCAGGGA CAACAGCAAC GAATAACCTT TAGTGAGACA GGCGCAGAGG 1000 AAGAGGAGGA GAATGACCAG CTGCCACTGG TGCATTGTGT GGTCGCCCGC 1050 TCGGAGAAAT CTGGTCTCCT CCAGGAACCC CTTGCGGGAG GCCCCCTGTC 1100 GAACCTCTCA GAAAACACGT GTGACAACTT TTGCTTGGTG CTTGAGGCAA 1150 TACAGGAGAA CCTGAGGCAG TACAAGGAGA TCCGAGATGA ACTCAACATG 1200 TAG

SEO ID NO: 2 (Human CatSper 3 Protein Sequence) MRDNEKAWWQ QWTSHTGLEG WGGTQEDRMG FGGAVAALRG RPSPLQSTIH 0050 ESYGRPEEQV LINRQEITNK ADAWDMQEFI THMYIKQLLR HPAFQLLLAL 0100 0150 LLVINAITIA LRTNSYLDQK HYELFSTIDD IVLTILLCEV LLGWLNGFWI 0200 FWKDGWNILN FIIVFILLLR FFINEINIPS INYTLRALRL VHVCMAVEPL ARIIRVILQS VPDMANIMVL ILFFMLVFSV FGVTLFGAFV PKHFQNIQVA 0250 0300 LYTLFICITQ DGWVDIYSDF QTEKREYAME IGGAIYFTIF ITIGAFIGIN 0350 LFVIVVTTNL EOMMKAGEOG OOORITFSET GAEEEEENDQ LPLVHCVVAR SEKSGLLQEP LAGGPLSNLS ENTCDNFCLV LEAIQENLRQ YKEIRDELNM 0400

SEQ ID NO: 3 (Murine CatSper 3 cDNA) ATGTCTGAAA AACACAAGTG GTGGCAGCAG GTGGAGAACA TCGACATCAC 0050 ACACCTGGGC CCTAAGAGAA AAGCCTATGA ACTCCTGGGT CGGCATGAGG 0100 AGCAAGTGCT CATCAACCGC AGAGATGTCA TGGAGAAGAA GGATGCCTGG 0150 GATGTACAGG AATTCATCAC TCAAATGTAT ATCAAGCAGT TGCTCCGCCA 0200 TCCGGCCTTC CAGCTGCTGC TGGCCTTTCT GCTGCTGTCC AACGCCATCA 0250 0300 CCATTGCCCT TCGCACCAAC TCTTATCTCG GTCAGAAACA CTACGAGCTA TTCTCGACCA TAGATGACAT TGTGTTGACG ATCCTTATCT GCGAGGTTCT 0350 GCTTGGTTGG CTTAACGGCT TCTGGATTTT CTGGAAGGAT GGCTGGAATA 0400 TCCTCAACTT CGCAATTGTC TTTATCTTGT TTATGGGGTT CTTCATAAAA 0450 CAACTTGACA TGGTTGCCAT CACCTACCCT CTCAGGGTGC TCCGGCTGGT 0500 GCATGTGTGT ATGGCGGTGG AACCCCTGGC CAGAATCATC AAGGTTATCC 0550 TGCAGTCGAT GCCAGACTTG GCCAATGTCA TGGCTCTCAT CCTCTTCTTC 0600 ATGCTGGTAT TCTCTGTGTT TGGGGTCACG CTCTTCGGTG CATTTGTGCC 0650 CAAGCATTTC CAGAACATGG GGGTTGCCCT GTACACGCTC TTCATCTGCA 0700 TCACTCAGGA TGGATGGCTG GACATCTACA CTGACTTCCA GATGGATGAA 0750 AGAGAGTACG CGATGGAGGT CGGGGGCGCC ATCTACTTTG CCGTCTTTAT 0800 CACCCTCGGT GCCTTCATTG GTCTCAACTT GTTCGTCGTC GTGGTGACCA 0850 CAAACCTGGA ACAAATGATG AAGACCGGCG AGGAAGAGGG ACACCTGAAC 0900 ATAAAGTITA CTGAGACAGA AGAGGATGAG GACTGGACCG ACGAGCTGCC 0950 ACTGGTGCAT TGTACAGAGG CCCGCAAGGA TACTTCCACT GTCCCCAAGG 1000 AACCACTGGT TGGGGGCCCC CTGAGTAACC TCACAGAAAA GACCTGCGAT 1050 AACTTCTGCT TGGTGCTTGA AGCAATACAG GAGAACTTGA TGGAGTACAA 1100 AGAGATCCGA GAGGAACTCA ACATGATCGT GGAGGAAGTG TCCTCCATCC 1150

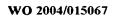


10/523479

GGTTCAACCA GGAGCAGCAA AATGTGATCC TACACAAGTA TACCTCCAAA	1200
AGCGCCACCT TCCTAAGCGA GCCCCCAGAA GGGGCTAACA AGCAAGACTT	1250
GATCACTGCG CTGGTCAGCA GGGAAAAGGT GTCTGATTCT AACATAAACA	1300
TGGTTAACAA ACACAAGTTC AGCCACTGA	1329

SEQ ID NO: 4 (Murine CatSper 3 Protein Sequence)	2252
MSEKHKWWQQ VENIDITHLG PKRKAYELLG RHEEQVLINR RDVMEKKDAW	0050
DVQEFITQMY IKQLLRHPAF QLLLAFLLLS NAITIALRTN SYLGQKHYEL	0100
FSTIDDIVLT ILICEVLLGW LNGFWIFWKD GWNILNFAIV FILFMGFFIK	0150
QLDMVAITYP LRVLRLVHVC MAVEPLARII KVILQSMPDL ANVMALILFF	0200
MLVFSVFGVT LFGAFVPKHF QNMGVALYTL FICITQDGWL DIYTDFQMDE	0250
REYAMEVGGA IYFAVFITLG AFIGLNLFVV VVTTNLEQMM KTGEEEGHLN	0300
IKFTETEEDE DWTDELPLVH CTEARKDTST VPKEPLVGGP LSNLTEKTCD	0350
NFCLVLEAIQ ENLMEYKEIR EELNMIVEEV SSIRFNQEQQ NVILHKYTSK	0400
SATFLSEPPE GANKQDLITA LVSREKVSDS NINMVNKHKF SH*	0442

SEQ ID NO: 5 (hCatSper3 5' flanking sequence containing basal promoter region)	
GGGCTGCCGG GGGTAGGAGG TGGGGATAAA CAACAGGGCG TGGAGCTCAG	0050
ACAGAAACCC TCTGTGCTTT CCACCCTGCC TGCAGCCCAG CCCTGCTCAA	0100
GCTGGGAGTC CCCTCCATGG AGACACATCA CCTGCAGCCA CCCCCACACA	0150
GCGCAGCCCA CGGACCTCCT TTGGCTCTCT GACAGGTGCT GGGCTGGAGT	0200
TGGGAGCTGG GCTGGGGGCT GGGGTGGGCA CATCCTCATC CTGCTCTTCC	0250
CTCCCACAGA CAGCAGTGAA GAGGCACTGG AAGGAATGGT ACGGGGGCTG	0300
AGGCAGGGTG GCGTGTCCCT CCTAGGCCAG CCACAGCCCC TGACCCAGGA	0350
ACAGTGGCGG AGCTCTTTCA TGCGGCGCAA CCGAGACCCT CAGCTCAATG	0400
AGCGAGTGCA CCGTGTGCGG GCGCTACAGA GCACACTCAA GGTCAGCTGG	0450
GGGGCTCTGG GCACAGCAAG GGACTAGGCT CTGGGCTTCA GGCTTTGGTT	0500
TGCGGCTGTC ACCTCCACCC TGGGCACCAG ACTCCAGACT CCAGACTCAG	0550
CTCCGGACCC TGGGCTTAGC AGCTGACAGC GGGCTCAGCT GTGGACTGGG	0600
CCAGGCTCTG GGTTCCGAGT GGGGATTTGA GTCTCACCTA GGCTCCTCGT	0650
GCCACGCTGG CCAGGTGCTG GCTTCCAGGC ACCGGACCTC CGGAGTGAAG	0700
TCTGGCCTCG GGCTCTGCCC ACTTCCCTGG GTGATCATGG TCCCTTAGCC	0750
CCTCCTCTCC ACACAGGCAA AGCTGCAGGA GCTGCAGGTC CTAGAAGAAG	0800
TGCTGGGTGA CCCTGAGCTG ACAGGAGAGA AGTTCCGCCA GTGGAAGGAG	0850
CAGAACCGGG AGCTGTACTC AGAGGGCCTG GGGGCCTGGG GAGTGGCACA	0900
GGCTGAAGGC AGCTCCCACA TCTTGACCTC TGACTCCACA GAACAGTCCC	0950
CCCACTCCCT GCCCTCTGAC CCTGAAGAGC ACTCCCATCT CTGCCCCCTG	1000
ACCTCAGAGA GCAGCCTCCG ACCTCCTGAC CTCTGACCCT GGCCAGCACT	1050
CTAGCTCCTG ACCTTTGACC CGAGGGCCAC CTCAACCCCA GCTTCTGACG	1100
TGTCCAGGAC AGAGCATCCC TGGATTCTGT TCAGGGTGGG AAGTAGTACT	1150
GCTAGTCATG GTCTCACCCC GAGCTGACCC CTCTGCCTGG GCTTTGTGCC	1200
ACCCTCTCCC TTGCCAAAGA AGAAACTCTC CCCCCAAATC CTCCAACCTC	1250
TGGGGCCACA GCCCTGCCCC TCCAGTTCCT TGGCAGTTCT CCCCCAAACC	1300
AGGTCTGTAC AGGTGTTCTT TATTTTACAT GAGGGCTACT TTCCAACCAA	1350
ATAAAGTCAA TTTTTCTAAG AATGAGTCTA CATGTAACTT TACTTCCATA	1400
TTCGAATTGG AAATCTGCCC CCCTGTGGGG ACTGGGGTGA GTGCTCTTGG	1450
CCAGAGGGTG GGTGGCAGAC CCTTCGTGCA GCCCGTTGGC CTGGGCTCTG	1500
TACCCGAGCT CCAAGCCTGC CAGGATGGTG GGGGATGACC CATGGCTAAT	1550
GAGGGCTCCG ACTCATGTCC ACCTCTCCCC AGCTCTTTGA AGGCTAATGG	1600
TGATCTCCTA CCCCATTCCC GGGGGGCACA CAATGAGAAA CTTCCACTTT	1650
GTAGATGGGG AAATGCACTT TGCACGGAAA GGTGGTGGGG ACAGTCCTGG	1700
AGACTGGGCT GGTAGGACAG GGCAGCTGGT GGGGAAGGGT GCAGGTTGAG	1750
GTCTGCCCTG GGAAGGCCCT GGGGAAAACA CTTCTCTCCT TCACTCCTCA	1800
TTCCAGCCTC ACCTCCACCT CCTGGATCCA AGGCAGGGAC ATGTCCCTGT	1850
GACTCCATTC AGGCTGCACG GGAAATCTGA CCTGCTCCCA TCAGCCTCTG	1900
ACTTCCAACC CCAGCCCAGC ATCCCCACAG CATCCCCAGA CTTCCTCTGT	1950
GGGATGCGGA GGAGGCCAA TGGGAGGAGC TTCTCTCCAG GTTGGAATTC	2000
CTCAGTAGAA TGCAGACGGC TGGAGGTCAC AGAGGCCTCT GTGATATCAC	2050
CACGAGGGG AGTGAGACCA CTTGGAGTG	2079





PCT/US2003/024432

10/523479

DT01 Rec'd PCT/PTC .0 4 FEB 2005

SEQ ID NO: 6 (hCatSper3 5' UTR)	•
AAGATTCTTT GAGGAGAAGG AAGAGACTGA GCAAA	\C

SEQ ID NO: 7 (hCatSper3 3' UTR)	
GGGAGGGTAC TGGGGCTGCC CCCAAGTCAT GTGAGTCAAG GCTGGGCGGA	0050
GCGTCAGAGT CTTCTGGCCT TCACGCCCTC ACCATTTATA AGGCAGAGCC	0100
TGGGCCCAC AGAGGTCCCC CACCCTATTG GTGGAGGAAC TGGAATCCAG	0150
ACTCCAGGTT CCTTCCATCT CACACAAGGG CACAGCTCGG CCTGGGTCTC	0200
TGTCAGGGCT GCGTGGGAGA GCGAAGCGGG GGTGACGCCA GGGAAGAGGT	0250
GGGAGGGCTG CTTCCCTCCC CTGAGGCCTT CTGAAAGGCA CTCACTGCTC	0300
CACCCCAGG ATTGTGGAGG AGGTGCGTGC AATCCGCTTC AACCAGGAGC	0350
AGGAGTCAGA GGTGTTGAAC AGGCGCTCGT CGACGAGCGG GTCGTTGGAG	0400
ACTACGTCAT CCAAGGACAT CCGCCAGATG TCTCAACAGC AAGACTTGCT	0450
CAGTGCGCTC GTTAGCATGG AAAAGGTG	0478